

RAW SEQUENCE LISTING

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Application Serial Number: 10/671,995 B
Source: IFW16
Date Processed by STIC: 05/05/2006

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IFW16

RAW SEQUENCE LISTING

DATE: 05/05/2006

PATENT APPLICATION: US/10/671,995B

TIME: 08:59:47

Input Set : A:\BIO1100.ST25.txt

Output Set: N:\CRF4\05052006\J671995B.raw

GOLD

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3 <110> APPLICANT: BIOHESION, INC.
4   WOODBURY, Richard
5   deVOS, Theo
6   IRANI, Meher
8 <120> TITLE OF INVENTION: RECOMBINANT FUSION PROTEINS WITH HIGH AFFINITY BINDING TO
AND APPLICATIONS THEREOF
9   AND APPLICATIONS THEREOF
11 <130> FILE REFERENCE: BIO1100
13 <140> CURRENT APPLICATION NUMBER: US 10/671,995B
14 <141> CURRENT FILING DATE: 2003-09-26
16 <160> NUMBER OF SEQ ID NOS: 30
18 <170> SOFTWARE: PatentIn version 3.3
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45 ttc tat gag atc tta cat tta cct aac tta aac gaa gaa caa cga aac      261
46 Phe Tyr Glu Ile Leu His Leu Pro Asn Leu Asn Glu Glu Gln Arg Asn
47           35              40              45
49 gcc ttc atc caa agt tta aaa gat gac cca agc caa agc gct aac ctt      309
50 Ala Phe Ile Gln Ser Leu Lys Asp Asp Pro Ser Gln Ser Ala Asn Leu
51 50              55              60              65
53 tta gca gaa gct aaa aag cta aat gat gct cag gcg ccg aaa gta gac      357
54 Leu Ala Glu Ala Lys Lys Leu Asn Asp Ala Gln Ala Pro Lys Val Asp
55           70              75              80
57 aac aaa ttc aac aaa gaa caa caa aac gcg ttc tat gag atc tta cat      405
58 Asn Lys Phe Asn Lys Glu Gln Gln Asn Ala Phe Tyr Glu Ile Leu His
59           85              90              95
61 tta cct aac tta aac gaa gaa caa cga aac gcc ttc atc caa agt tta      453
62 Leu Pro Asn Leu Asn Glu Glu Gln Arg Asn Ala Phe Ile Gln Ser Leu

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66	Lys Asp Asp Pro Ser Gln Ser Ala Asn Leu Leu Ala Glu Ala Lys Lys			
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69	cta aat gat gct cag gcg ccg aaa gta gac gcg aat tcg agc tct ggt	549		
70	Leu Asn Asp Ala Gln Ala Pro Lys Val Asp Ala Asn Ser Ser Ser Gly			
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73	agt ggc aat ggt cat atg cat gga aaa act cag gca acc agc ggg act	597		
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78	Ile Gln Ser Met His Gly Lys Thr Gln Ala Thr Ser Gly Thr Ile Gln			
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81	agc atg cat gga aaa act cag gca acc agc ggg act atc cag agc atg	693		
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85	cat gga aaa act cag gca acc agc ggg act atc cag agc atg cat gga	741		
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87	195	200	205	
89	aaa act cag gca acc agc ggg act atc cag agc atg cat gga aaa act	789		
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93	cag gca acc agc ggg act atc cag agc atg cat gga aaa att cag gca	837		
94	Gln Ala Thr Ser Gly Thr Ile Gln Ser Met His Gly Lys Ile Gln Ala			
95	230	235	240	
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98	Thr Ser Gly Thr Ile Gln Ser Met His Ala Leu Ser Leu Glu Gly Pro			
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147	tggcgcgcat	tgcgcccagc	gccatctgat	cggtggcaac	cagcatcgca	gtgggaacga	2325
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276 50 55 60
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288 100 105 110
291 Leu Lys Asp Asp Pro Ser Gln Ser Ala Asn Leu Leu Ala Glu Ala Lys
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296 130 135 140
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300 145 150 155 160
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307 Gln Ser Met His Gly Lys Thr Gln Ala Thr Ser Gly Thr Ile Gln Ser
308 180 185 190
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312 195 200 205
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316 210 215 220
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350                                     5      10      15
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354      20      25      30
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362 50      55      60      65
364 tac gac tcc gct ccg gct acc gac ggt tcc gga acc gct ctg ggt tgg      357
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